

FIG. 1

APPROVED	0.3 FIG.
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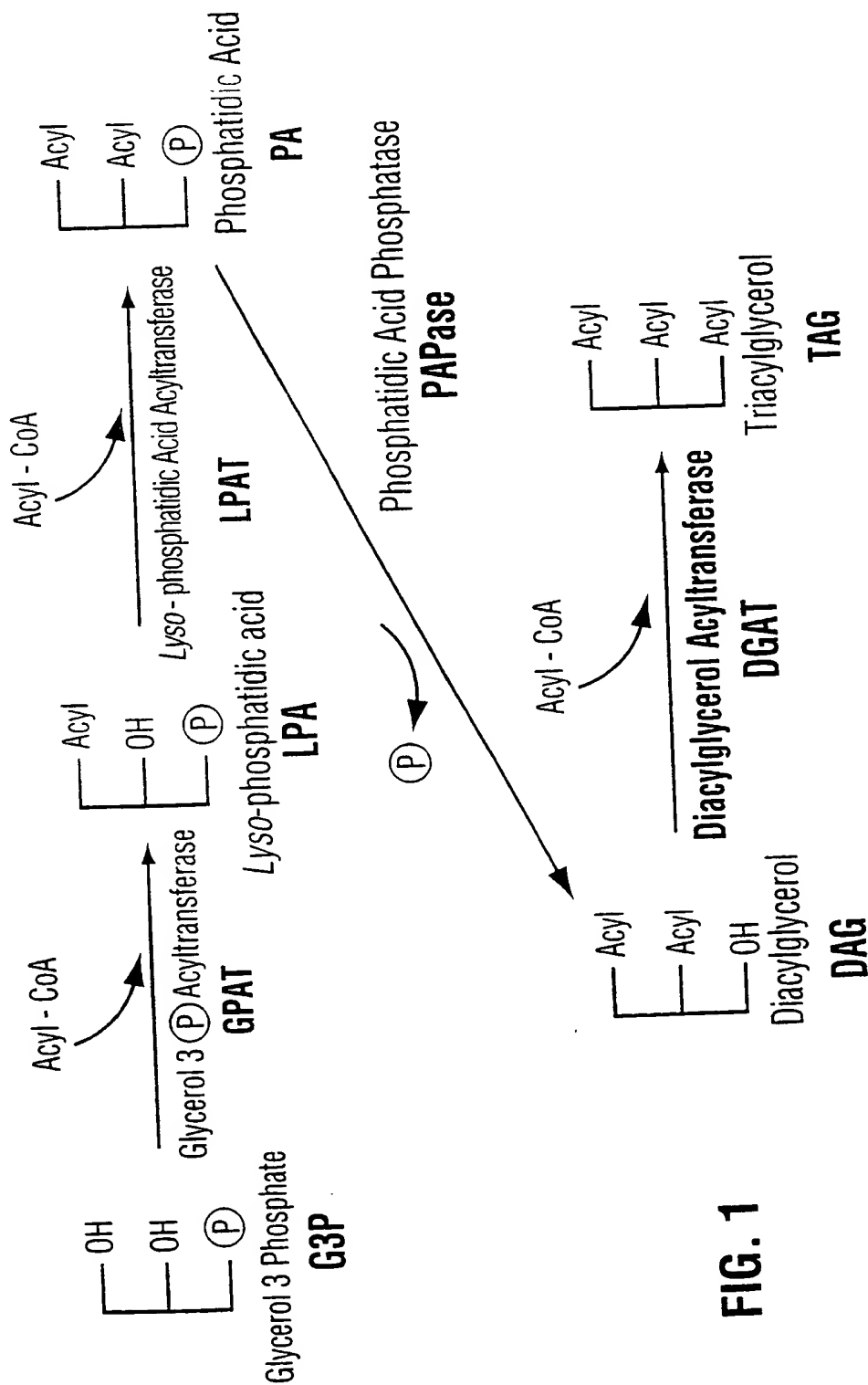


FIG. 1

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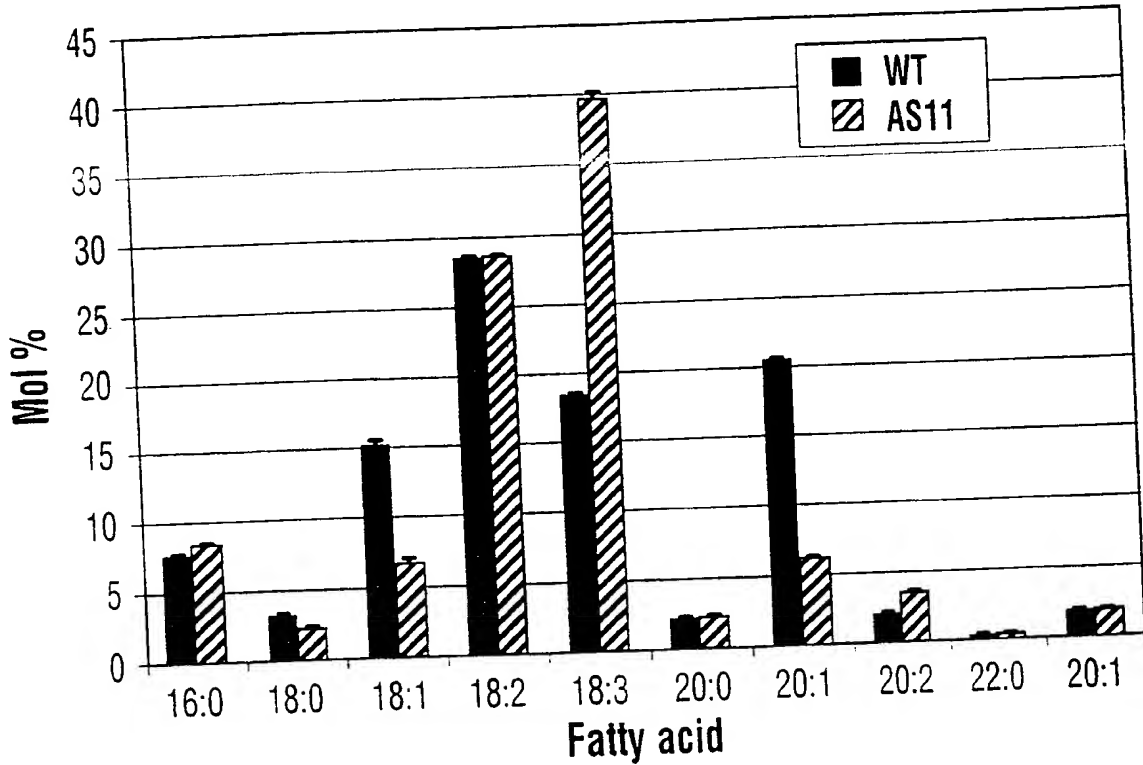


FIG. 2

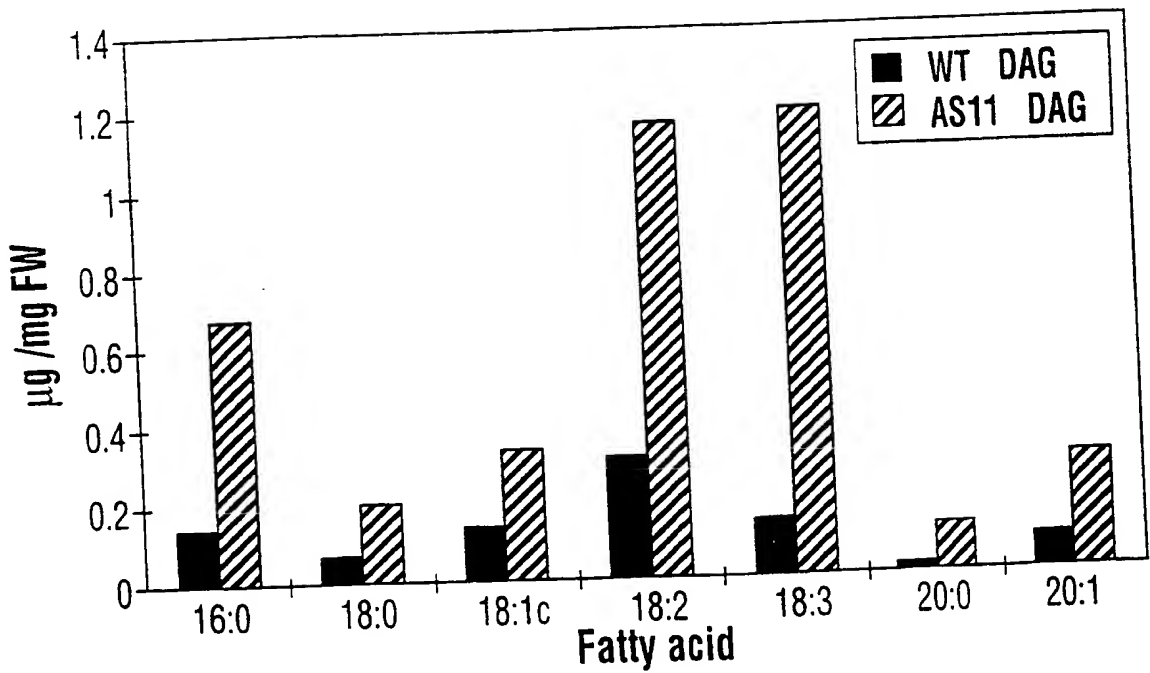


FIG. 3

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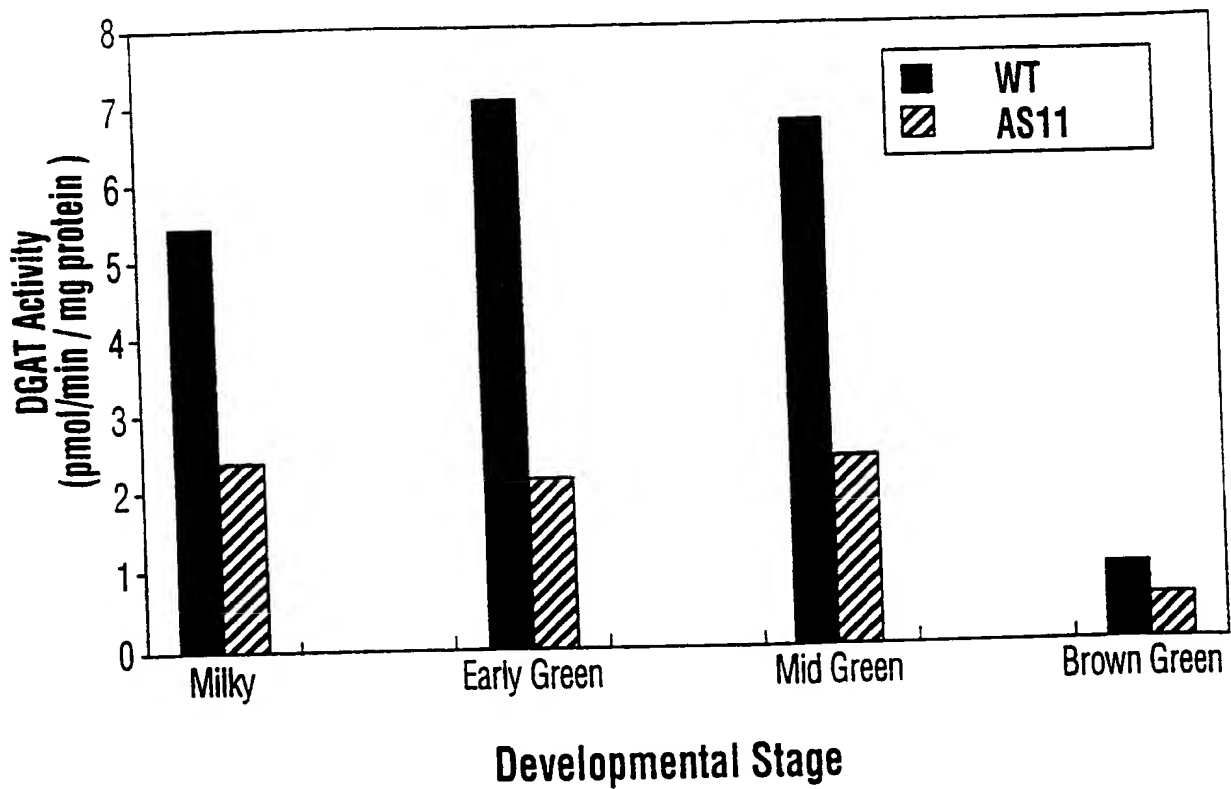


FIG. 4

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(a)

AtTAG1 MAILDSAGVTTVTENGGEFVDLDRLRRRKSRSDSSNGLLLSGSDNNSPS 50
 MDGAT MDRGGAGSSR.....RRRTGSRVSVQG.....GSGPKVEE 31
 HARGP1 MGDR...GSSR.....RRRTGSRPSSHG.....GGPAAAE 28

AtTAG1 DEVGAPADVRDRIDSVVNDDAQGTANLAGDNNGGGDNNGGGRGCGEGRGN 100
 MDGAT DEVRDAAVSPDL.....GAGGDAPAPAPAPAHTRDKDGRTSVCECY.. 72
 HARGP1 DEVRDAAGPDV.....GAAGDAPAPAP.....NKDGDAGVCSGH.. 63

AtTAG1 ADATETYPSPVPAHRRARESPLSSDIIFKQSHAGIFNL CVVVLIAVNR 150
 MDGATWDLRC.....HRLQDSLSSDSGFSNYR.GILNWCVVMLILSNARL 112
 HARGP1WELRC.....HRLQDSLSSDSGFSNYR.GILNWCVVMLILSNARL 103

AtTAG1 IIFENLMKYGWLIRTDWFSSRSLED...WPLFMC.CISLSIFPEAAFTYE 196
 MDGAT FIFENLMKYGILVDP.IQVVSFLKDPYSWPA.PCVIIASNIFVMAAFQTE 160
 HARGP1 FIFENLMKYGILVDP.IQVVSFLKDPHSWPA.PCLVIAANVEFAVAAFQTE 151

AtTAG1 KLVLOKYSEEPVVIIFLHIIITMTEVLYPVYVTLRCDSAFLSGVTMLLTC 246
 MDGAT KRLAVGALTEQMGLLLHVNLATITCEPAAVALLVESITPVGSVFALASY 210
 HARGP1 KRLAVGALTEQAGLLLVANLATITCEPAAVLLVESITPVGSLALMAH 201

AtTAG1 .IVWLKLVSYAHTSY.....DIRSLANAADR.....ANPEVSYVSL.. 282
 MDGAT SIMELKLYSYRDVNLWCRQRRVKAKAVSTGKKVSGAAQQAUSYDPNLTY 260
 HARGP1 TIFELKLYSYRDVNSWC..RRARAKAASAGKKASSAAPHTVSYDPNLTY 249

AtTAG1 KSLAYFVAVPTLCYQPSYPRACIRKGVVARQFAKLVIFTGFMGFITIEQV 332
 MDGAT RDLIYFIFAPPTLCYELNEPRSPRIRKRELLRRVLEMEFFTQLOVGLIQOW 310
 HARGP1 RDLIYFIFAPPTLCYELNEPRSPRIRKRELLRRILEMEFFTQLOVGLIQOW 299

AtTAG1 INPIVRNSKHPLKGDILYA..IERVLKLSVPNLYVWLCMFYCFHFLWLNI 380
 MDGAT MVPTTONSMKPEK.DMDYSRIIERELKLA VPNHLIWLIFFYWFFHSCNLA 359
 HARGP1 MVPTTONSMKPEK.DMDYSRIIERELKLA VPNHLIWLIFFYWLFHSCNLA 348

AtTAG1 LAELFCFGDREFYKDWNAKSVGDYWRMWNMPVHKWVVRHIVFPCLRSKI 430
 MDGAT VAELFCFGDREFYRDWNAESVITYEWNWNIPVHKWCTRHFKPMLRHGS 409
 HARGP1 VAELMFCFGDREFYRDWNNSESVITYEWNWNIPVHKWCTRHFKPMLRRGS 398

AtTAG1 PKTBAIIIAFLVSAVFHELCHAVPCREFKLWAFLGIMFOVPLV.FITNXL 479
 MDGAT SKWMARTGVFLTSAFFHEYLSVPLRMFRLWAF TAMMAQPLAWIVGREF 459
 HARGP1 SKWMARTGVFLASAFFHEYLSVPLRMFRLWAF TGMMAQPLAWFVGREF 448

AtTAG1 QERFGSTVGNMIFWFIFCFIFGQPMCVLEYYHD..EMNRKGSMS.. 520
 MDGAT QGNYGNA.....VWVTL.IIGQPVAVLMYVHDYYVENVYDAPVG.V 498
 HARGP1 QGNYGNA.....VWLSL.IIGQPEAVLMYVHDYYVENVYEAFAEA 488

FIG. 5(a)

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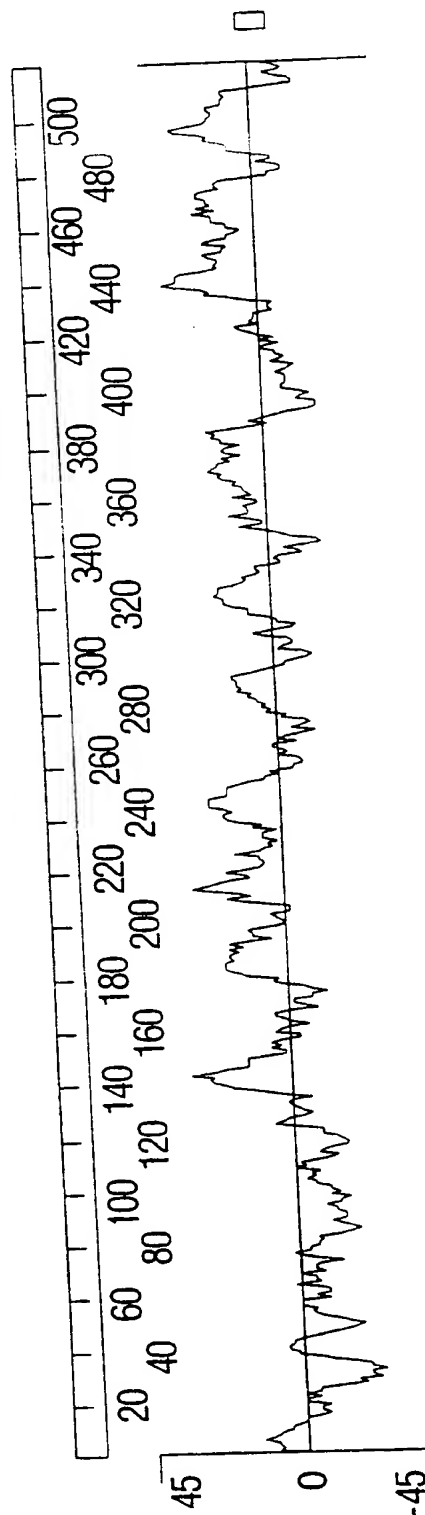


FIG. 5(b)

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RT LF FL YS SL GS



FIG. 6(a)

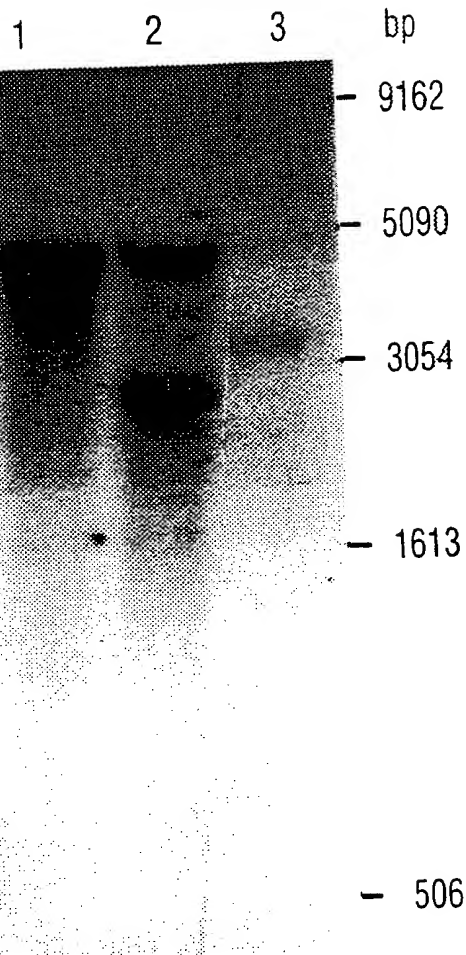


FIG. 6(b)

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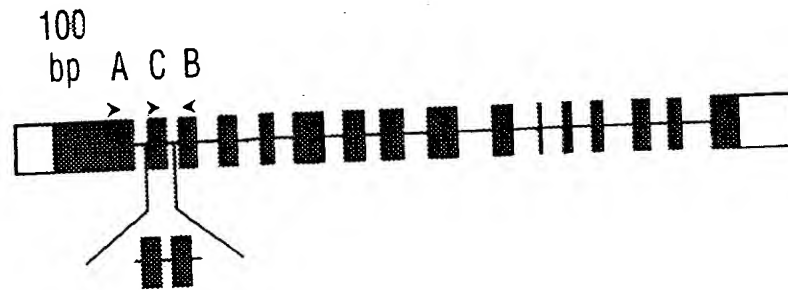


FIG. 7(a)

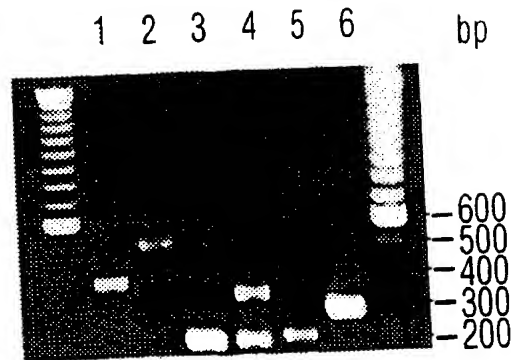


FIG. 7(b)

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APPROVED	FIG. 8
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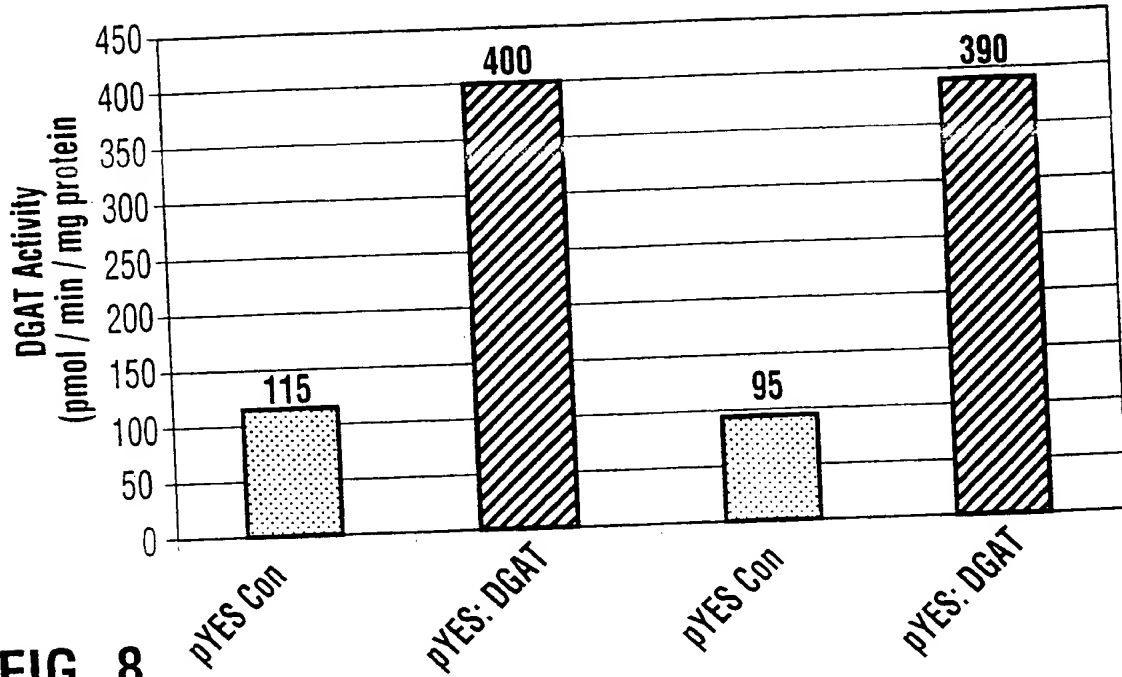


FIG. 8

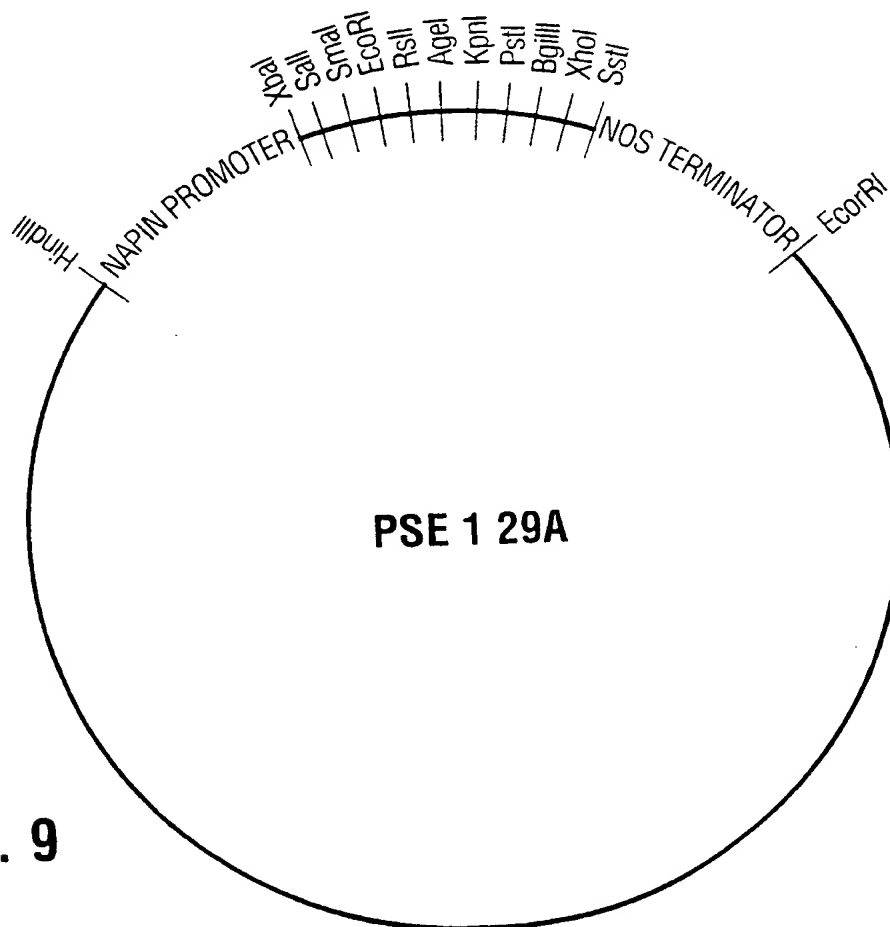


FIG. 9

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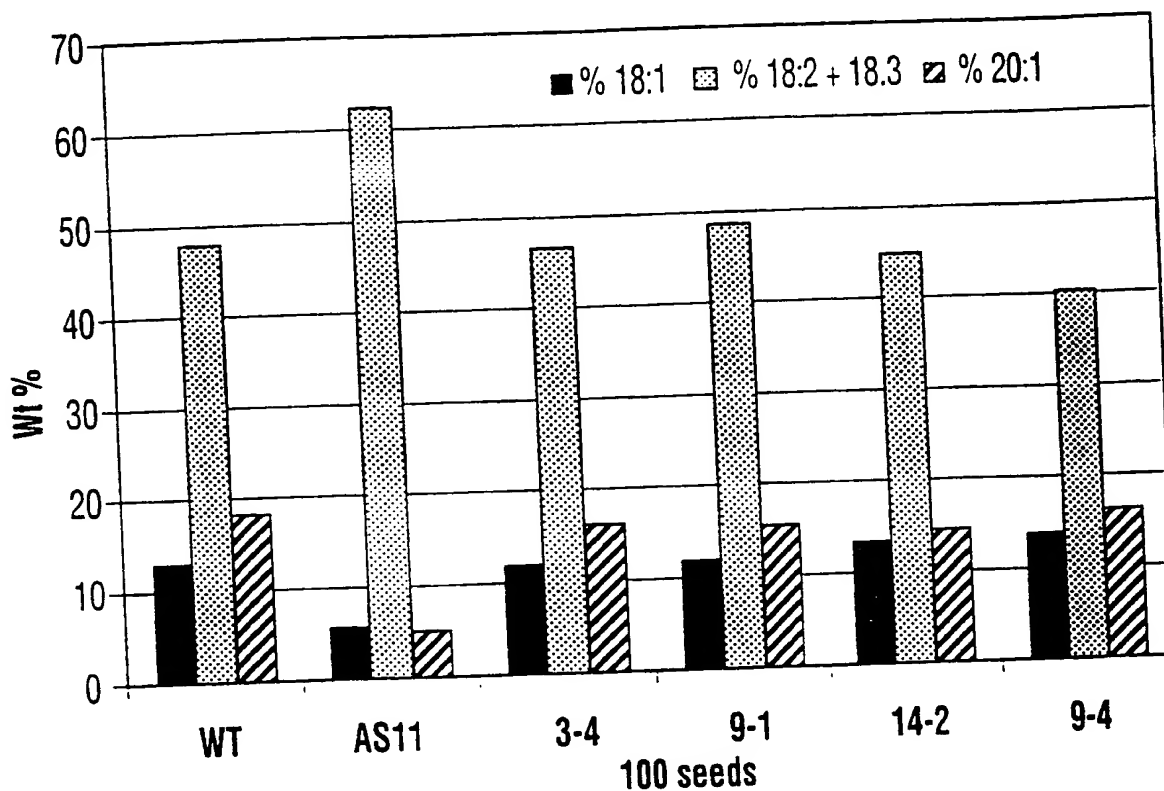


FIG. 10

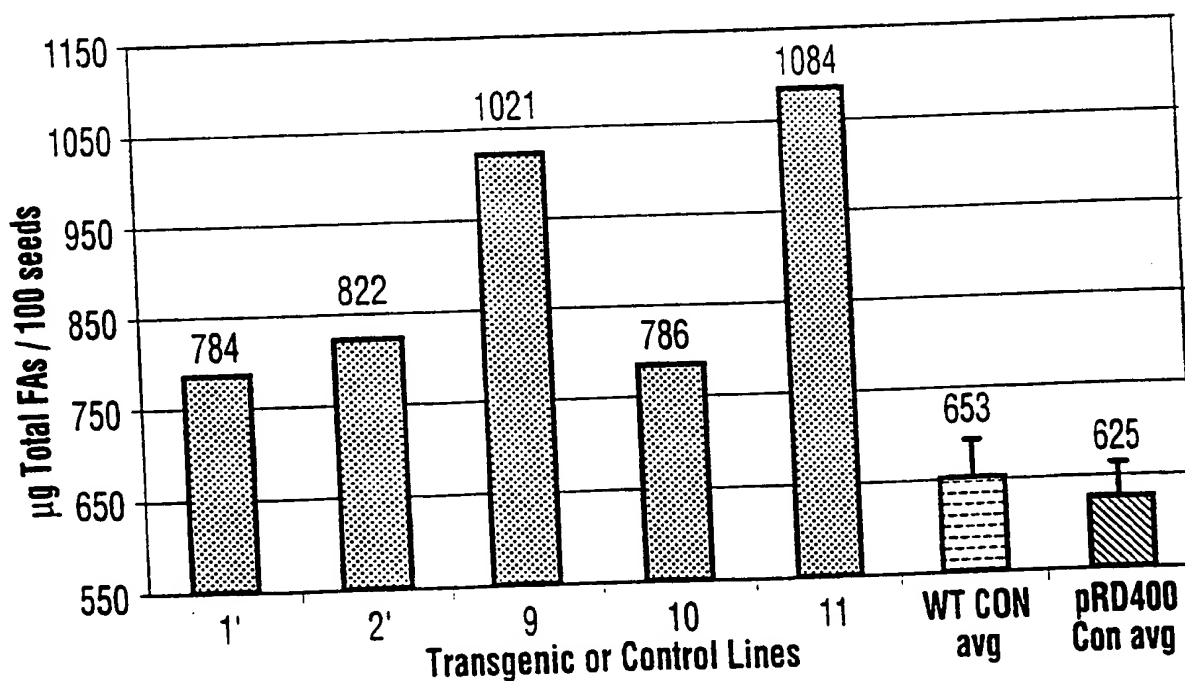


FIG. 11

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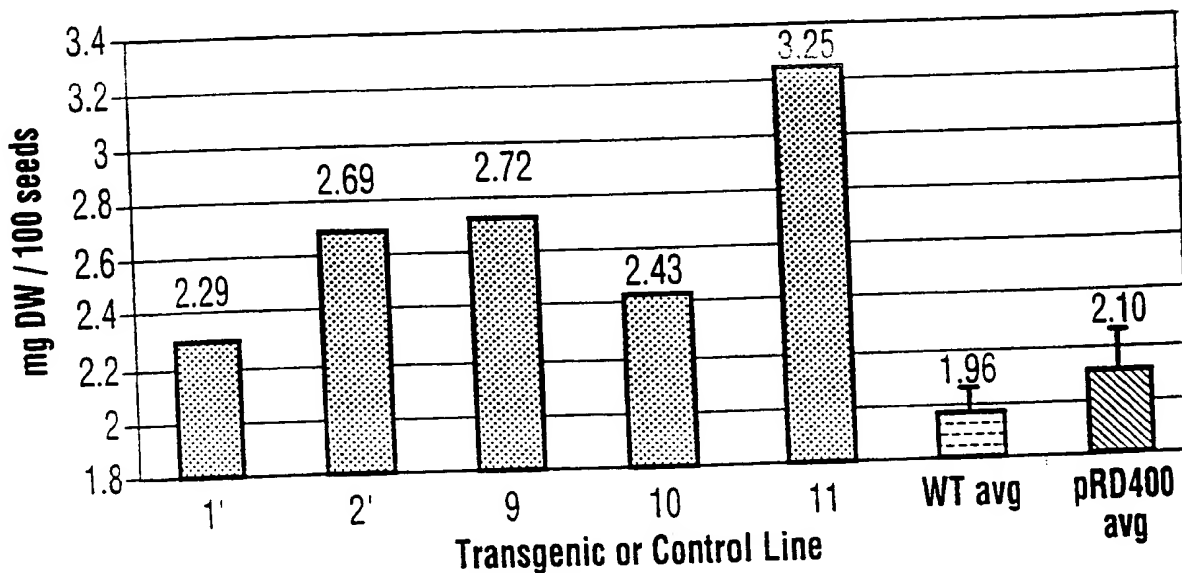


FIG. 12

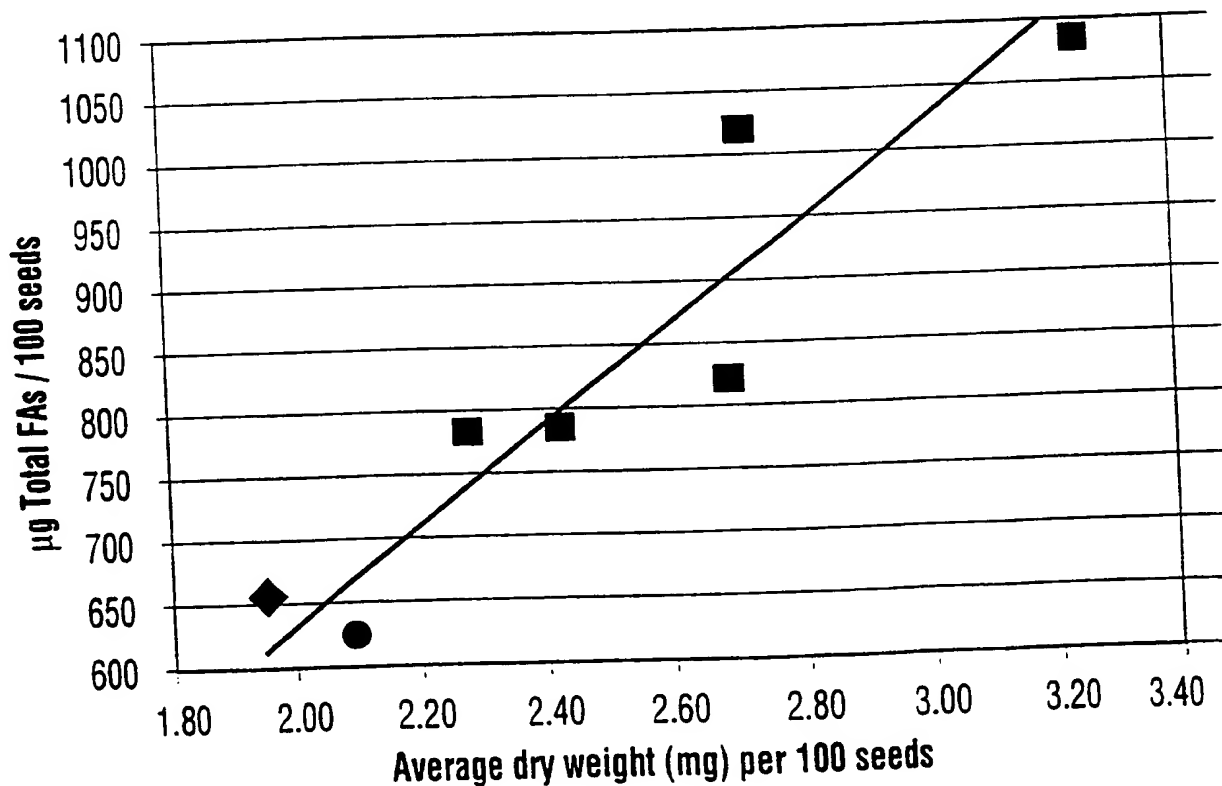


FIG. 13